PCT OPERATIONS

FACSIMILE TRANSMISSION COVER SHEET

DATE: April 19, 2001
TO: Natalie Berzta
TELEPHONE: (202) 721-5491 FAX NO.: (202) 530-1055
FROM: Barbara Comphell
TELEPHONE: <u>703 305-363/</u> FAX NO.: 703-308-4785 OR 703-305-3230
MESSAGE: Erran Repart that was

NUMBER OF PAGES (INCLUDING THIS PAGE)

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FCH&S D.C.

92025301055

TE67

PGS. SENT RESULT

ST. TIME

CONNECTION ID

SUBADDRESS CONNECTION TEL

TX/RX NO

TRANSMISSION OK



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable

Application Serial Number:

Source:

Date Processed by STIC:

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER: 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE

2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0 The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 = 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual

Property Organization (WIPO) Standard ST.25

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence

sion 3.0 can be down loaded from the USPTO website at the following address: http://www.aisprogov/web/offices/pac/checkers

< 1107 - list applicant 1207 & give invertion to be 1407 & U.S. serial number (application) Does Not Comply
1417 & U.S. filing date
1607 & Rotal number 1 securior 09/674,973 The submitted file SEQUENCE LISTING COMMON FOR ALL SEQUENCES: (a portion storen as a sample) is invalid SEQUENCE TYPE: Poptide SEQUENCE UNIT: Amino Acid describe organism: Attificial Sequence, of Scientific hame (benus/species) RHPSWPWTRCLRMRPPRS My Unknown or SEQUENCE ID NO: 2 SEQUENCE LENGTH: 31 amino acids Artificial Sequence G T R A G P G P G A S G C V H Q E A E R V S Q A H R G R T G Aplain in 62237 20 section SEQUENCE ID NO: 3 SEQUENCE LENGTH: 32 amino acids GGTRAGPGPGASGCVHQEAERVSQAHRGRT 30 20 les Sequence Rules, do not use one-letter amero acids. Use thee-letter amino acids and rumby them under every 5 amino acids. DNOT use TAB codes between ameno acid numbers. Use space characters. Please: I) consult new Sequera Rules 2) consult sample Sequera Listing, attacked, for valid format.

Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least on in or Xaa.

Ø 003

<110> smith, John

Smith, Jane

<120> Example of a Sequence Listing

<130> 01-00001

<140> US 08/999,999

<141> 1998-02-28

<150> EP 91000000

<151> 1997-12-31

Please consult

ME EAST

<160> 2

<170> PatentIn ver. 2.0

<210> 1

<211> 403

<212> DNA

<213> Paramecium aurelia

<220>

<221> CDS

<222> 341..394

<30.0>

<301> Doe, Richard

<302> Isolation and Characterization of a Gene Encoding a

Protease from Paramecium sp.

<303> Journal of Fictional Genes

Zana Zana

<305× 4

<306> I - 7

<307> 1988-06-20

<400> 1

ctactctact ctactctcat ctactatett ctttggatet ctgagtetge ctgagtggta 60

ctettgagte etggagatet eteeteteae atgtgategt egagaetgae egatagateg 120

ctgactgact ctgagatagt cgageccgta cgagacccgt cgagggtgac agagagtggg 180

cgcgtgcgcg cagagcgccg cgccggtgcg cgcgcgagtg cgcggtgggc cgcgcgaggg 240

etttegegge ageggeggeg ettteeggeg egegeeegte egeceetaga eetgagaggt 300

cttetettee etectettea etagagaggt etatatatae atg gtt tea atg tte 355

Met Val Ser Met Phe

age tig tet tie and tig cet gga tit tigt tig tit git tigtitigete

403

Ser Leu Ser Phe Lys Trp Pro Gly Phe Cys Leu Phe Val

خذ

<210> 2

<211> 18

<212> PRT

<213> Paramecium aurclia

<400> 2

200 B

USPTO/PCT OPEP

DESC GUE EUL THA LE FT TURELLE

1

10

15

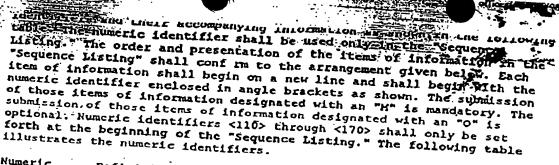
Phe Val

ed: May 22, 1998,
A. Lehman,
ant Secretary of Commerce and
alssioner of Patents and Trademarks,
oc. 98–14194 Filed 5–29–98; 8:45 aml
a cooc assa-140

800

OSPTO/PCT OPEP

ULY 606 601 LA1 16.61 10 62.40



•	Marchie 1	dentifiers.	ruc rollowing table
Numeri Identii	Dossars		
	ret.	Comments and Format	Mandatory (H) or Optional (O)
<110>	Applicant	Preferably max. of 10 names; one name per line; preferable format; Surname, Other Names, and/or	H →
<120>	Title of	Initials.	
	Invention	•	м
<130>	File Reference	.	
	٠	reference	M when filed prior to assignment of appl. number
<140>	Current Application Number	a- Specify as: US 07/999,999 or PCT/US96/99999	M, if available
<141>	Current filing	Specify as: yyyy-mm-d	ld M, if available
<150>	Prior Application		M, if applicable include priority documents under 35 USC 119 and
<151>	Prior Application	n Specify as: yyyy-mm-dd	
<160>	Number of SEQ ID NOs		м
<170>	Software	Name of software used to create the Sequence Listing	o
<210>	SEQ ID NO:#:	Response shall be an integer representing the SEQ ID NO shown	м
<211>		Respond with an integer expressing the number of bases or amino acid residues	
			-

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Whether presented sequence molecule is DNA, RNA, or PRT (protein). If a nucleotide sequence contains both DNA . and RNA fragments, the type shall be "DNA." in acdition, the combined DNA/ .RNA molecule shall be further described in

<213>

Organism

Scientific name, i.e. Genus/species, Unknown or Artificial Sequence. In addition, the "Unknown" or "Artificial Scquence" organisms shall be further described in the <220> to <223> feature section.

the`<220> to <223} feature

<220>

Feature

Leave blank after <220>. <221-223> provide for a description of points of biological significance in the sequence.

M, under the following conditions: if "n,"
"Xaa," or a modified or unusual L-amino acid or modified base was used in a sequence; if ORGAN-ISM is "Artificial Sequence" or "Unknown"; if molecule is combined DNA/RNA.

М

<221>

Name/Key

Provide appropriate identifier for feature, preferably from WIPO Standard ST.25 (1998), Appendix 2, Tables 5 and 6

M, under the following conditions:
if "n," "Xaa," or
a modified or unusual L-amino
acid or modified
base was used in
a sequence

<222>

Location

Specify location within sequence; where appropriate state number of first and last bases/amino acids

M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified

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in feature

a gequer

<223>		a Sequent					
<300>	Other Information	Other relevant information; four lines maximum		M, under the fol-			
10005	Publication Information	Leave blank after <300>		0			
<301>	Authors	Preferably max of ten named authors of publication; specify one name per line; preferable format: Surname, Other Names and/or Initials		o			
<302>	Title						
<303>	Journal	•	0			•	
<304>	Volume		. 0				
<305>	Issue		0				
<306>	Pages ·		0				
<307>	Date	Journal date on which data published; specify as yyyy-mm-dd, MMM-yyyy or Season-yyyy	0				
<308>	Database Accession Number	Accession number assigned by data-base including database name	o				
<309>	Database Entry Date	Date of entry in database; specify as yyyy-mm-dd or MMM-yyyy	0				
<310>		Document number; for patent-type citations only. Specify as, for example; US 07/999,999	o				

of 14

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Document filing date, for patenttype citations only; specify as yyyy-ma-dd

<312>

Publication Date Document publication date, for Patent-type citations only; specify as yyyy-mm-dd

<313> Relevant

Residues

FROM (position) TO (position)

K

<400> Sequence

SEQ ID NO should follow the numeric identifier and should appear on the line preceding the actual sequence

5. Section 1.824 is revised to read as follows:

1.824 Form and format for nucleotide and/or amino acid sequence submissions in computer readable form.

- (a) The computer readable form required by following specifications: 1.821(c) shall meet the
- (1) The computer readable form shall contain a single "Sequence Listing" as either a diskette, series of diskettes, or other permissible media outlined in paragraph (c) of this section.
- (2) The "Sequence Listing" in paragraph (a) (1) of this section shall be submitted in American Standard Code for Information Interchange (ASCII)
- (3) The computer readable form may be created by any means, such as word processors, nucleotide/amino acid sequence editors or other custom Computer programs; however, it shall conform to all specifications
- (4) File compression is acceptable when using diskette media, so long as the compressed file is in a self-extracting format that will decompress on one of the systems described in paragraph (b) of this section.
- (5) Page numbering shall not appear within the computer readable form version of the "Sequence Listing" file.
- (6) All computer readable forms shall have a label permanently affixed thereto on which has been hand-printed or typed: the name of the applicant, the title of the invention, the date on which the data were recorded on the computer readable form, the operating system used, a reference number, and an application serial number and filing date, if
- (b) Computer readable form submissions must meet these format requirements:
- (1) Computer: IBM PC/XT/AT, or compatibles, or Apple Macintosh;
- (2) Operating System: MS-DOS, Unix or Macintosh;

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-hancdi Pilm